

UNITED STATES PATENT AND TRADEMARK OFFICE
CERTIFICATE OF CORRECTION

PATENT NO. : 6,872,546 B1
DATED : March 29, 2005
INVENTOR(S) : Hastings et al.

Page 1 of 60

It is certified that error appears in the above-identified patent and that said Letters Patent is hereby corrected as shown below:

Title page.

Item [56], **References Cited**, OTHER PUBLICATIONS, fourth reference should read:
-- Lee et al., "A novel secretory Tumor Necrosis Factor-inducible Protein (TSG-6) is a member of the family of Hyaluronate Binding Proteins, closely related to the Adhesion Receptor CD44" Jan. 1992, vol. 116, No. 2, pp. 545-557.* --.

Column 315.

Starting at line 36, before claims, insert the attached sequence listing.

Column 317.

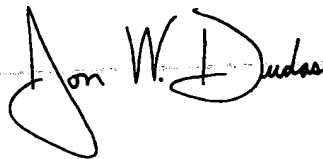
Lines 11-19, delete the second subsets (a)-(g).

Column 318.

Line 9, after "claim 33," insert -- wherein said --.

Signed and Sealed this

Fourth Day of April, 2006

A handwritten signature in black ink, appearing to read "Jon W. Dudas". The signature is stylized with a large, looped initial "J" and a cursive "Dudas".

JON W. DUDAS
Director of the United States Patent and Trademark Office

SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 37

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 Met Met Asp
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 5 10 15

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 Glu Asp Thr Arg Thr Gln Gln Thr Arg Arg Trp Trp Thr Leu Ala Gly
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cag gag atc acc gtc acc ttt aac caa ttc acg aaa tac tcc tac aag 418
 Gln Glu Ile Thr Val Thr Phe Asn Gln Phe Thr Lys Tyr Ser Tyr Lys
 70 75 80

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Leu Ala Ser Thr Glu Ala Phe Ser Arg Phe Glu Thr Ile Leu Glu Asn	
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Val Asn Gly Ile Leu His Phe Ile Asp Arg Val Leu Leu Pro Pro Glu	
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Pro Gln Leu Ser Ala Ala Gln Gln Leu Gly Phe His Leu Cys Leu Met	
1830 1835 1840	
ggc tgg ctg gcc aat ggc tcc act gcc cac cct gtg gtt ttc cct gtg	5746
Gly Trp Leu Ala Asn Gly Ser Thr Ala His Pro Val Val Phe Pro Val	
1845 1850 1855	

gcg gac tgt ggc aat ggt cgg gtg ggc ata gtc agc ctg ggt gcc cgc	5794
Ala Asp Cys Gly Asn Gly Arg Val Gly Ile Val Ser Leu Gly Ala Arg	
1860 1865 1870 1875	
aag aac ctc tca gaa cgc tgg gat gcc tac tgc ttc cgt gtg caa gat	5842
Lys Asn Leu Ser Glu Arg Trp Asp Ala Tyr Cys Phe Arg Val Gln Asp	
1880 1885 1890	
gtg gcc tgc cga tgc cga aat ggc ttc gtg ggt gac ggg atc agc acg	5890
Val Ala Cys Arg Cys Arg Asn Gly Phe Val Gly Asp Gly Ile Ser Thr	
1895 1900 1905	
tgc aat ggg aag ctg ctg gat gtg ctg gct gcc act gcc aac ttc tcc	5938
Cys Asn Gly Lys Leu Leu Asp Val Leu Ala Ala Thr Ala Asn Phe Ser	
1910 1915 1920	
acc ttc tat ggg atg cta ttg ggc tat gcc aat gcc acc cag cgg ggt	5986
Thr Phe Tyr Gly Met Leu Leu Gly Tyr Ala Asn Ala Thr Gln Arg Gly	
1925 1930 1935	
ctc gac ttc ctg gac ttc ctg gat gat gag ctc acg tat aag aca ctc	6034
Leu Asp Phe Leu Asp Phe Leu Asp Asp Glu Leu Thr Tyr Lys Thr Leu	
1940 1945 1950 1955	
ttc gtc cct gtc aat gaa ggc ttt gtg gac aac atg acg ctg agt ggc	6082
Phe Val Pro Val Asn Glu Gly Phe Val Asp Asn Met Thr Leu Ser Gly	
1960 1965 1970	
cca aac ttg gag ctg cat gcc tcc aac gcc acc ctc cta agt gcc aac	6130
Pro Asn Leu Glu Leu His Ala Ser Asn Ala Thr Leu Leu Ser Ala Asn	
1975 1980 1985	
gcc agc cag ggg aag ttg ctt ccg gcc cac tca ggc ctc agc ctc atc	6178
Ala Ser Gln Gly Lys Leu Leu Pro Ala His Ser Gly Leu Ser Leu Ile	
1990 1995 2000	
atc agt gac gca ggc cct gac aac agt tcc tgg gcc cct gtg gcc cca	6226
Ile Ser Asp Ala Gly Pro Asp Asn Ser Ser Trp Ala Pro Val Ala Pro	
2005 2010 2015	

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 Ala Ser Leu Ala Gln Gln Leu Cys Arg Gln His Ile Ile Ala Gly Gln
 35 40 45
 His Ile Leu Glu Asp Thr Arg Thr Gln Gln Thr Arg Arg Trp Trp Thr
 50 55 60

Leu	Ala	Gly	Gln	Glu	Ile	Thr	Val	Thr	Phe	Asn	Gln	Phe	Thr	Lys	Tyr	65	70	75	80
Ser	Tyr	Lys	Tyr	Lys	Asp	Gln	Pro	Gln	Gln	Thr	Phe	Asn	Ile	Tyr	Lys	85	90	95	
Ala	Asn	Asn	Ile	Ala	Ala	Asn	Gly	Val	Phe	His	Val	Val	Thr	Gly	Leu	100	105	110	
Arg	Trp	Gln	Ala	Pro	Ser	Gly	Thr	Pro	Gly	Asp	Pro	Lys	Arg	Thr	Ile	115	120	125	
Gly	Gln	Ile	Leu	Ala	Ser	Thr	Glu	Ala	Phe	Ser	Arg	Phe	Glu	Thr	Ile	130	135	140	
Leu	Glu	Asn	Cys	Gly	Leu	Pro	Ser	Ile	Leu	Asp	Gly	Pro	Gly	Pro	Phe	145	150	155	160
Thr	Val	Phe	Ala	Pro	Ser	Asn	Glu	Ala	Val	Asp	Ser	Leu	Arg	Asp	Gly	165	170	175	
Arg	Leu	Ile	Tyr	Leu	Phe	Thr	Ala	Gly	Leu	Ser	Lys	Leu	Gln	Glu	Leu	180	185	190	
Val	Arg	Tyr	His	Ile	Tyr	Asn	His	Gly	Gln	Leu	Thr	Val	Glu	Lys	Leu	195	200	205	
Ile	Ser	Lys	Gly	Arg	Ile	Leu	Thr	Met	Ala	Asn	Gln	Val	Leu	Ala	Val	210	215	220	
Asn	Ile	Ser	Glu	Glu	Gly	Arg	Ile	Leu	Leu	Gly	Pro	Glu	Gly	Val	Pro	225	230	235	240
Leu	Gln	Arg	Val	Asp	Val	Met	Ala	Ala	Asn	Gly	Val	Ile	His	Met	Leu	245	250	255	
Asp	Gly	Ile	Leu	Leu	Pro	Pro	Thr	Ile	Leu	Pro	Ile	Leu	Pro	Lys	His	260	265	270	
Cys	Ser	Glu	Glu	Gln	His	Lys	Ile	Val	Ala	Gly	Ser	Cys	Val	Asp	Cys	275	280	285	

Gln	Ala	Leu	Asn	Thr	Ser	Thr	Cys	Pro	Pro	Asn	Ser	Val	Lys	Leu	Asp	290	295	300	
Ile	Phe	Pro	Lys	Glu	Cys	Val	Tyr	Ile	His	Asp	Pro	Thr	Gly	Leu	Asn	305	310	315	320
Val	Leu	Lys	Lys	Gly	Cys	Ala	Ser	Tyr	Cys	Asn	Gln	Thr	Ile	Met	Glu	325	330	335	
Gln	Gly	Cys	Cys	Lys	Gly	Phe	Phe	Gly	Pro	Asp	Cys	Thr	Gln	Cys	Pro	340	345	350	
Gly	Gly	Phe	Ser	Asn	Pro	Cys	Tyr	Gly	Lys	Gly	Asn	Cys	Ser	Asp	Gly	355	360	365	
Ile	Gln	Gly	Asn	Gly	Ala	Cys	Leu	Cys	Phe	Pro	Asp	Tyr	Lys	Gly	Ile	370	375	380	
Ala	Cys	His	Ile	Cys	Ser	Asn	Pro	Asn	Lys	His	Gly	Glu	Gln	Cys	Gln	385	390	395	400
Glu	Asp	Cys	Gly	Cys	Val	His	Gly	Leu	Cys	Asp	Asn	Arg	Pro	Gly	Ser	405	410	415	
Gly	Gly	Val	Cys	Gln	Gln	Gly	Thr	Cys	Ala	Pro	Gly	Phe	Ser	Gly	Arg	420	425	430	
Phe	Cys	Asn	Glu	Ser	Met	Gly	Asp	Cys	Gly	Pro	Thr	Gly	Leu	Ala	Gln	435	440	445	
His	Cys	His	Leu	His	Ala	Arg	Cys	Val	Ser	Gln	Glu	Gly	Val	Ala	Arg	450	455	460	
Cys	Arg	Cys	Leu	Asp	Gly	Phe	Glu	Gly	Asp	Gly	Phe	Ser	Cys	Thr	Pro	465	470	475	480
Ser	Asn	Pro	Cys	Ser	His	Pro	Asp	Arg	Gly	Gly	Cys	Ser	Glu	Asn	Ala	485	490	495	
Glu	Cys	Val	Pro	Gly	Ser	Leu	Gly	Thr	His	His	Cys	Thr	Cys	His	Lys	500	505	510	

Gly Trp Ser Gly Asp Gly Arg Val Cys Val Ala Ile Asp Glu Cys Glu
515 520 525

Leu Asp Val Arg Gly Gly Cys His Thr Asp Ala Leu Cys Ser Tyr Val
530 535 540

Gly Pro Gly Gln Ser Arg Cys Thr Cys Lys Leu Gly Phe Ala Gly Asp
545 550 555 560

Gly Tyr Gln Cys Ser Pro Ile Asp Pro Cys Arg Ala Gly Asn Gly Gly
565 570 575

Cys His Gly Leu Glu Leu Glu Ala Asn Ala His Phe Ser Ile Phe Tyr
580 585 590

Gln Trp Leu Lys Ser Ala Gly Ile Thr Leu Pro Ala Asp Arg Arg Val
595 600 605

Thr Ala Leu Val Pro Ser Glu Ala Ala Val Arg Gln Leu Ser Pro Glu
610 615 620

Asp Arg Ala Phe Trp Leu Gln Pro Arg Thr Leu Pro Asn Leu Val Arg
625 630 635 640

Ala His Phe Leu Gln Gly Ala Leu Phe Glu Glu Glu Leu Ala Arg Leu
645 650 655

Gly Gly Gln Glu Val Ala Thr Leu Asn Pro Thr Thr Arg Trp Glu Ile
660 665 670

Arg Asn Ile Ser Gly Arg Val Trp Val Gln Asn Ala Ser Val Asp Val
675 680 685

Ala Asp Leu Leu Ala Thr Asn Gly Val Leu His Ile Leu Ser Gln Val
690 695 700

Leu Leu Pro Pro Arg Gly Asp Val Pro Gly Gly Gln Gly Leu Leu Gln
705 710 715 720

Gln Leu Asp Leu Val Pro Ala Phe Ser Leu Phe Arg Glu Leu Leu Gln
725 730 735

His	His	Gly	Leu	Val	Pro	Gln	Ile	Glu	Ala	Ala	Thr	Ala	Tyr	Thr	Ile	740	745	750
Phe	Val	Pro	Thr	Asn	Arg	Ser	Leu	Glu	Ala	Gln	Gly	Asn	Ser	Ser	His	755	760	765
Leu	Asp	Ala	Asp	Thr	Val	Arg	His	His	Val	Val	Leu	Gly	Glu	Ala	Leu	770	775	780
Ser	Met	Glu	Thr	Leu	Arg	Lys	Gly	Gly	His	Arg	Asn	Ser	Leu	Leu	Gly	785	790	795
Pro	Ala	His	Trp	Ile	Val	Phe	Tyr	Asn	His	Ser	Gly	Gln	Pro	Glu	Val	805	810	815
Asn	His	Val	Pro	Leu	Glu	Gly	Pro	Met	Leu	Glu	Ala	Pro	Gly	Arg	Ser	820	825	830
Leu	Ile	Gly	Leu	Ser	Gly	Val	Leu	Thr	Val	Gly	Ser	Ser	Arg	Cys	Leu	835	840	845
His	Ser	His	Ala	Glu	Ala	Leu	Arg	Glu	Lys	Cys	Val	Asn	Cys	Thr	Arg	850	855	860
Arg	Phe	Arg	Cys	Thr	Gln	Gly	Phe	Gln	Leu	Gln	Asp	Thr	Pro	Arg	Lys	865	870	875
Ser	Cys	Val	Tyr	Arg	Ser	Gly	Phe	Ser	Phe	Ser	Arg	Gly	Cys	Ser	Tyr	885	890	895
Thr	Cys	Ala	Lys	Lys	Ile	Gln	Val	Pro	Asp	Cys	Cys	Pro	Gly	Phe	Phe	900	905	910
Gly	Thr	Leu	Cys	Glu	Pro	Cys	Pro	Gly	Gly	Leu	Gly	Gly	Val	Cys	Ser	915	920	925
Gly	His	Gly	Gln	Cys	Gln	Asp	Arg	Phe	Leu	Gly	Ser	Gly	Glu	Cys	His	930	935	940
Cys	His	Glu	Gly	Phe	His	Gly	Thr	Ala	Cys	Glu	Val	Cys	Glu	Leu	Gly	945	950	955
																		960

Arg Tyr Gly Pro Asn Cys Thr Gly Val Cys Asp Cys Ala His Gly Leu
 965 970 975

Cys Gln Glu Gly Leu Gln Gly Asp Gly Ser Cys Val Cys Asn Val Gly
 980 985 990

Trp Gln Gly Leu Arg Cys Asp Gln Lys Ile Thr Ser Pro Gln Cys Pro
 995 1000 1005

Arg Lys Cys Asp Pro Asn Ala Asn Cys Val Gln Asp Ser Ala Gly Ala
 1010 1015 1020

Ser Thr Cys Ala Cys Ala Ala Gly Tyr Ser Gly Asn Gly Ile Phe Cys
 1025 1030 1035 1040

Ser Glu Val Asp Pro Cys Ala His Gly His Gly Gly Cys Ser Pro His
 1045 1050 1055

Ala Asn Cys Thr Lys Val Ala Pro Gly Gln Arg Thr Cys Thr Cys Gln
 1060 1065 1070

Asp Gly Tyr Met Gly Asp Gly Glu Leu Cys Gln Glu Ile Asn Ser Cys
 1075 1080 1085

Leu Ile His His Gly Gly Cys His Ile His Ala Glu Cys Ile Pro Thr
 1090 1095 1100

Gly Pro Gln Gln Val Ser Cys Ser Cys Arg Glu Gly Tyr Ser Gly Asp
 1105 1110 1115 1120

Gly Ile Arg Thr Cys Glu Leu Leu Asp Pro Cys Ser Lys Asn Asn Gly
 1125 1130 1135

Gly Cys Ser Pro Tyr Ala Thr Cys Lys Ser Thr Gly Asp Gly Gln Arg
 1140 1145 1150

Thr Cys Thr Cys Asp Thr Ala His Thr Val Gly Asp Gly Leu Thr Cys
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Arg Ala Arg Val Gly Leu Glu Leu Leu Arg Asp Lys His Ala Ser Phe
 1170 1175 1180

Phe Ser Leu Arg Leu Leu Glu Tyr Lys Glu Leu Lys Gly Asp Gly Pro
 1185 1190 1195 1200

Phe Thr Ile Phe Val Pro His Ala Asp Leu Met Ser Asn Leu Ser Gln
 1205 1210 1215

Asp Glu Leu Ala Arg Ile Arg Ala His Arg Gln Leu Val Phe Arg Tyr
 1220 1225 1230

His Val Val Gly Cys Arg Arg Leu Arg Ser Glu Asp Leu Leu Glu Gln
 1235 1240 1245

Gly Tyr Ala Thr Ala Leu Ser Gly His Pro Leu Arg Phe Ser Glu Arg
 1250 1255 1260

Glu Gly Ser Ile Tyr Leu Asn Asp Phe Ala Arg Val Val Ser Ser Asp
 1265 1270 1275 1280

His Glu Ala Val Asn Gly Ile Leu His Phe Ile Asp Arg Val Leu Leu
 1285 1290 1295

Pro Pro Glu Ala Leu His Trp Glu Pro Asp Asp Ala Pro Ile Pro Arg
 1300 1305 1310

Arg Asn Val Thr Ala Ala Ala Gln Gly Phe Gly Tyr Lys Ile Phe Ser
 1315 1320 1325

Gly Leu Leu Lys Val Ala Gly Leu Leu Pro Leu Leu Arg Glu Ala Ser
 1330 1335 1340

His Arg Pro Phe Thr Met Leu Trp Pro Thr Asp Ala Ala Phe Arg Ala
 1345 1350 1355 1360

Leu Pro Pro Asp Arg Gln Ala Trp Leu Tyr His Glu Asp His Arg Asp
 1365 1370 1375

Lys Leu Ala Ala Ile Leu Arg Gly His Met Ile Arg Asn Val Glu Ala
 1380 1385 1390

Leu Ala Ser Asp Leu Pro Asn Leu Gly Pro Leu Arg Thr Met His Gly
 1395 1400 1405

Thr	Pro	Ile	Ser	Phe	Ser	Cys	Ser	Arg	Thr	Arg	Pro	Gly	Glu	Leu	Met	
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Val	Gly	Glu	Asp	Asp	Ala	Arg	Ile	Val	Gln	Arg	His	Leu	Pro	Phe	Glu	
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Gly	Ala	Arg	Cys	Asp	His	Phe	Glu	Thr	Arg	Pro	Leu	Arg	Leu	Asn	Thr	
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Cys	Ser	Ile	Cys	Gly	Leu	Glu	Pro	Pro	Cys	Pro	Glu	Gly	Ser	Gln	Glu	
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Gln	Gly	Ser	Pro	Glu	Ala	Cys	Trp	Arg	Phe	Tyr	Pro	Lys	Phe	Trp	Thr	
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Ser	Pro	Pro	Leu	His	Ser	Leu	Gly	Leu	Arg	Ser	Val	Trp	Val	His	Pro	
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Ser	Leu	Trp	Gly	Arg	Pro	Gln	Gly	Leu	Gly	Arg	Gly	Cys	His	Arg	Asn	
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Cys	Val	Thr	Thr	Thr	Trp	Lys	Pro	Ser	Cys	Cys	Pro	Gly	His	Tyr	Gly	
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Ser	Glu	Cys	Gln	Ala	Cys	Pro	Gly	Gly	Pro	Ser	Ser	Pro	Cys	Ser	Asp	
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Arg	Gly	Val	Cys	Met	Asp	Gly	Met	Ser	Gly	Ser	Gly	Gln	Cys	Leu	Cys	
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Phe	Gly	Pro	His	Cys	Gln	Ala	Cys	Arg	Cys	Thr	Val	His	Gly	Arg	Cys	
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Asp	Glu	Gly	Leu	Gly	Gly	Ser	Gly	Ser	Cys	Phe	Cys	Asp	Glu	Gly	Trp	
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Thr Gly Pro	Arg Cys Glu Val	Gln Leu Glu Leu	Gln Pro Val Cys Thr
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Pro Pro Cys Ala Pro	Glu Ala Val Cys Arg Ala	Gly Asn Ser Cys Glu	
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Cys Ser Leu Gly Tyr	Glu Gly Asp Gly Arg Val	Cys Thr Val Ala Asp	
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Leu Cys Gln Asp Gly	His Gly Gly Cys Ser	Glu His Ala Asn Cys Ser	
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Gln Val Gly Thr	Met Val Thr Cys Thr	Cys Leu Pro Asp Tyr Glu Gly	
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Asp Gly Trp Ser Cys Arg Ala Arg	Asn Pro Cys Thr Asp Gly His Arg		
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Gly Gly Cys Ser Glu His Ala	Asn Cys Leu Ser Thr Gly Leu Asn Thr		
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Arg Arg Cys Glu Cys His Ala Gly Tyr Val	Gly Asp Gly Leu Gln Cys		
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Leu Glu Glu Ser Glu Pro Pro Val Asp Arg	Cys Leu Gly Gln Pro Pro		
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Pro Cys His Ser Asp Ala Met Cys Thr	Asp Leu His Phe Gln Glu Lys		
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Arg Ala Gly Val Phe His Leu Gln Ala Thr Ser Gly Pro Tyr Gly Leu			
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Asn Phe Ser Glu Ala Glu Ala Ala Cys Glu Ala Gln Gly Ala Val Leu			
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Ala Ser Phe Pro Gln Leu Ser Ala Ala Gln Gln Leu Gly Phe His Leu			
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Cys Leu Met Gly Trp Leu Ala Asn Gly Ser Thr Ala His Pro Val Val			
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Val	Gln	Asp	Val	Ala	Cys	Arg	Cys	Arg	Asn	Gly	Phe	Val	Gly	Asp	Gly				
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Ile	Ser	Thr	Cys	Asn	Gly	Lys	Leu	Leu	Asp	Val	Leu	Ala	Ala	Thr	Ala				
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Asn	Phe	Ser	Thr	Phe	Tyr	Gly	Met	Leu	Leu	Gly	Tyr	Ala	Asn	Ala	Thr				
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Gln	Arg	Gly	Leu	Asp	Phe	Leu	Asp	Phe	Leu	Asp	Asp	Glu	Leu	Thr	Tyr				
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Lys	Thr	Leu	Phe	Val	Pro	Val	Asn	Glu	Gly	Phe	Val	Asp	Asn	Met	Thr				
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Leu	Ser	Gly	Pro	Asn	Leu	Glu	Leu	His	Ala	Ser	Asn	Ala	Thr	Leu	Leu				
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Ser	Ala	Asn	Ala	Ser	Gln	Gly	Lys	Leu	Leu	Pro	Ala	His	Ser	Gly	Leu				
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Ser	Leu	Ile	Ile	Ser	Asp	Ala	Gly	Pro	Asp	Asn	Ser	Ser	Trp	Ala	Pro				
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Val	Ala	Pro	Gly	Thr	Val	Val	Val	Ser	Arg	Ile	Ile	Val	Trp	Asp	Ile				
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Met	Ala	Phe	Asn	Gly	Ile	Ile	His	Ala	Leu	Ala	Ser	Pro	Leu	Leu	Ala				
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Pro	Pro	Gln	Pro	Gln	Ala	Val	Leu	Ala	Xaa	Glu	Ala	Pro	Pro	Val	Ala				
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Ala	Gly	Val	Gly	Ala	Val	Leu	Ala	Ala	Gly	Ala	Leu	Leu	Gly	Leu	Val				
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Ala Gly Ala Leu Tyr Leu Arg Ala Arg Gly Lys Pro Met Gly Phe Gly
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Trp Gln Glu Gly Thr Asn Pro Xaa Leu Xaa Xaa Val Pro Asn Pro Val
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Ala Ala Lys Ala Val Cys Gly Gly His Ala Thr Tyr Lys Ala Ala Arg
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Lys Gly His Val Cys Ala Ala Gly Trp Met Ala Lys Gly Arg Val Gly
 50 55 60

Tyr Val Lys Gly Asn Cys Gly Gly Lys Thr Gly Asp Tyr Gly Arg Asn
 65 70 75 80

Arg Ser Arg Trp Asp Ala Tyr Cys Tyr Asn His Ala Lys Cys Gly Gly
 85 90 95

Val Thr Asp Lys Arg Lys Ser Gly Asn Tyr Asp Asn Cys Tyr Trp His
 100 105 110

Arg Lys Tyr Gly Arg His Ser Asp Asp Asp Asp Gly Cys Ala Asp Tyr
 115 120 125

Val Tyr Asp Ser Tyr Asp Asp Val His Gly Val Gly Arg Tyr Cys Gly
 130 135 140

Asp Asp Asp Ser Thr Gly Asn Val Met Thr Lys Ser Asp Ala Ser Val
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Thr Ala Gly Gly Lys Tyr Val Ala Met Asp Val Ser Lys Ser Ser Gly
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Lys Asn Thr Ser Thr Thr Ser Thr Gly Asn Lys Asn Ala Gly Arg Ser
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His

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Asp Tyr Glu Gly Asp Gly Trp Ser Cys Arg Ala Arg Asn Pro Cys Thr	
10 15 20	
gat ggc cac cgc ggg ggc tgc agc gag cac gcc aac tgc ttg agc acc	150
Asp Gly His Arg Gly Gly Cys Ser Glu His Ala Asn Cys Leu Ser Thr	
25 30 35 40	
ggc ctg aac aca cgg cgc tgt gag tgc cac gca ggc tac gta ggc gat	198
Gly Leu Asn Thr Arg Arg Cys Glu Cys His Ala Gly Tyr Val Gly Asp	
45 50 55	
gga ctg cag tgt ctg gag gag tcg gaa cca cct gtg gac cgc tgc ttg	246
Gly Leu Gln Cys Leu Glu Glu Ser Glu Pro Pro Val Asp Arg Cys Leu	
60 65 70	
ggc cag cca ccg ccc tgc cac tca gat gcc atg tgc act gac ctg cac	294
Gly Gln Pro Pro Pro Cys His Ser Asp Ala Met Cys Thr Asp Leu His	
75 80 85	
ttc cag gag aaa cgg gct ggc gtt ttc cac ctc cag gcc acc agc ggc	342
Phe Gln Glu Lys Arg Ala Gly Val Phe His Leu Gln Ala Thr Ser Gly	
90 95 100	
cct tat ggt ctg aac ttt tcg gag gct gag gcg gca tgc gaa gca cag	390
Pro Tyr Gly Leu Asn Phe Ser Glu Ala Glu Ala Ala Cys Glu Ala Gln	
105 110 115 120	
gga gcc gtc ctt gct tca ttc cct cag ctc tct gct gcc cag cag ctg	438
Gly Ala Val Leu Ala Ser Phe Pro Gln Leu Ser Ala Ala Gln Gln Leu	
125 130 135	
ggc ttc cac ctg tgc ctc atg ggc tgg ctg gcc aat ggc tcc act gcc	486
Gly Phe His Leu Cys Leu Met Gly Trp Leu Ala Asn Gly Ser Thr Ala	
140 145 150	
cac cct gtg gtt ttc cct gtg gcg gac tgt ggc aat ggt cgg gtg ggc	534
His Pro Val Val Phe Pro Val Ala Asp Cys Gly Asn Gly Arg Val Gly	
155 160 165	

ata gtc agc ctg ggt gcc cgc aag aac ctc tca gaa cgc tgg gat gcc	582
Ile Val Ser Leu Gly Ala Arg Lys Asn Leu Ser Glu Arg Trp Asp Ala	
170 175 180	
tac tgc ttc cgt gtg caa gat gtg gcc tgc cga tgc cga aat ggc ttc	630
Tyr Cys Phe Arg Val Gln Asp Val Ala Cys Arg Cys Arg Asn Gly Phe	
185 190 195 200	
gtg ggt gac ggg atc agc acg tgc aat ggg aag ctg ctg gat gtg ctg	678
Val Gly Asp Gly Ile Ser Thr Cys Asn Gly Lys Leu Leu Asp Val Leu	
205 210 215	
gct gcc act gcc aac ttc tcc acc ttc tat ggg atg cta ttg ggc tat	726
Ala Ala Thr Ala Asn Phe Ser Thr Phe Tyr Gly Met Leu Leu Gly Tyr	
220 225 230	
gcc aat gcc acc cag cgg ggt ctc gac ttc ctg gac ttc ctg gat gat	774
Ala Asn Ala Thr Gln Arg Gly Leu Asp Phe Leu Asp Phe Leu Asp Asp	
235 240 245	
gag ctc acg tat aag aca ctc ttc gtc cct gtc aat gaa ggc ttt gtg	822
Glu Leu Thr Tyr Lys Thr Leu Phe Val Pro Val Asn Glu Gly Phe Val	
250 255 260	
gac aac atg acg ctg agt ggc cca aac ttg gag ctg cat gcc tcc aac	870
Asp Asn Met Thr Leu Ser Gly Pro Asn Leu Glu Leu His Ala Ser Asn	
265 270 275 280	
gcc acc ctc cta agt gcc aac gcc agc cag ggg aag ttg ctt ccg gcc	918
Ala Thr Leu Leu Ser Ala Asn Ala Ser Gln Gly Lys Leu Leu Pro Ala	
285 290 295	
cac tca ggc ctc agc ctc atc atc agt gac gca ggc cct gac aac agt	966
His Ser Gly Leu Ser Leu Ile Ile Ser Asp Ala Gly Pro Asp Asn Ser	
300 305 310	
tcc tgg gcc cct gtg gcc cca ggg aca gtt gtg gtt agc cgt atc att	1014
Ser Trp Ala Pro Val Ala Pro Gly Thr Val Val Val Ser Arg Ile Ile	
315 320 325	

gtg tgg gac atc atg gcc ttc aat ggc atc atc cat gct ctg gcc agc	1062
Val Trp Asp Ile Met Ala Phe Asn Gly Ile Ile His Ala Leu Ala Ser	
330 335 340	
ccc ctc ctg gca ccc cca cag ccc cag gca gtg ctg gcg cnt gaa gcc	1110
Pro Leu Leu Ala Pro Pro Gln Pro Gln Ala Val Leu Ala Xaa Glu Ala	
345 350 355 360	
cca cct gtg gcg gca ggc gtg ggg gct gtg ctt gcc gct gga gca ctg	1158
Pro Pro Val Ala Ala Gly Val Gly Ala Val Leu Ala Ala Gly Ala Leu	
365 370 375	
ctt ggc ttg gtg gcc gga gct ctc tac ctc cgt gcc cga ggc aag ccc	1206
Leu Gly Leu Val Ala Gly Ala Leu Tyr Leu Arg Ala Arg Gly Lys Pro	
380 385 390	
atg ggc ttt ggc ttc tct gcc ttc cag gcg gaa gat gat gct gat gac	1254
Met Gly Phe Gly Phe Ser Ala Phe Gln Ala Glu Asp Asp Ala Asp Asp	
395 400 405	
gan ttc tca ccg tgg caa gaa ggg acc aac ccc acn ttg gtn tnt gtc	1302
Xaa Phe Ser Pro Trp Gln Glu Gly Thr Asn Pro Xaa Leu Xaa Xaa Val	
410 415 420	
ccc aac cct gtc ttt ggc agc gac acc ttt tgt gaa ccc ttc gat gac	1350
Pro Asn Pro Val Phe Gly Ser Asp Thr Phe Cys Glu Pro Phe Asp Asp	
425 430 435 440	
tca ctg ctg gag gag gac ttc cct gac acc cag agg atc ctc aca gtc	1398
Ser Leu Leu Glu Glu Asp Phe Pro Asp Thr Gln Arg Ile Leu Thr Val	
445 450 455	
aag tga cgaggctggg gctgaaagca gaagcatgca cagggaggag accantttta	1454
Lys	
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ccctcgag	1522

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 20 25 30
 Glu His Ala Asn Cys Leu Ser Thr Gly Leu Asn Thr Arg Arg Cys Glu
 35 40 45
 Cys His Ala Gly Tyr Val Gly Asp Gly Leu Gln Cys Leu Glu Glu Ser
 50 55 60

Glu	Pro	Pro	Val	Asp	Arg	Cys	Leu	Gly	Gln	Pro	Pro	Pro	Cys	His	Ser	65	70	75	80
Asp	Ala	Met	Cys	Thr	Asp	Leu	His	Phe	Gln	Glu	Lys	Arg	Ala	Gly	Val	85	90		95
Phe	His	Leu	Gln	Ala	Thr	Ser	Gly	Pro	Tyr	Gly	Leu	Asn	Phe	Ser	Glu	100	105		110
Ala	Glu	Ala	Ala	Cys	Glu	Ala	Gln	Gly	Ala	Val	Leu	Ala	Ser	Phe	Pro	115	120		125
Gln	Leu	Ser	Ala	Ala	Gln	Gln	Leu	Gly	Phe	His	Leu	Cys	Leu	Met	Gly	130	135		140
Trp	Leu	Ala	Asn	Gly	Ser	Thr	Ala	His	Pro	Val	Val	Phe	Pro	Val	Ala	145	150		155
Asp	Cys	Gly	Asn	Gly	Arg	Val	Gly	Ile	Val	Ser	Leu	Gly	Ala	Arg	Lys	165	170		175
Asn	Leu	Ser	Glu	Arg	Trp	Asp	Ala	Tyr	Cys	Phe	Arg	Val	Gln	Asp	Val	180	185		190
Ala	Cys	Arg	Cys	Arg	Asn	Gly	Phe	Val	Gly	Asp	Gly	Ile	Ser	Thr	Cys	195	200		205
Asn	Gly	Lys	Leu	Leu	Asp	Val	Leu	Ala	Ala	Thr	Ala	Asn	Phe	Ser	Thr	210	215		220
Phe	Tyr	Gly	Met	Leu	Leu	Gly	Tyr	Ala	Asn	Ala	Thr	Gln	Arg	Gly	Leu	225	230		235
Asp	Phe	Leu	Asp	Phe	Leu	Asp	Asp	Glu	Leu	Thr	Tyr	Lys	Thr	Leu	Phe	245	250		255
Val	Pro	Val	Asn	Glu	Gly	Phe	Val	Asp	Asn	Met	Thr	Leu	Ser	Gly	Pro	260	265		270
Asn	Leu	Glu	Leu	His	Ala	Ser	Asn	Ala	Thr	Leu	Leu	Ser	Ala	Asn	Ala	275	280		285

Ser Gln Gly Lys Leu Leu Pro Ala His Ser Gly Leu Ser Leu Ile Ile
 290 295 300

Ser Asp Ala Gly Pro Asp Asn Ser Ser Trp Ala Pro Val Ala Pro Gly
 305 310 315 320

Thr Val Val Val Ser Arg Ile Ile Val Trp Asp Ile Met Ala Phe Asn
 325 330 335

Gly Ile Ile His Ala Leu Ala Ser Pro Leu Leu Ala Pro Pro Gln Pro
 340 345 350

Gln Ala Val Leu Ala Xaa Glu Ala Pro Pro Val Ala Ala Gly Val Gly
 355 360 365

Ala Val Leu Ala Ala Gly Ala Leu Leu Gly Leu Val Ala Gly Ala Leu
 370 375 380

Tyr Leu Arg Ala Arg Gly Lys Pro Met Gly Phe Gly Phe Ser Ala Phe
 385 390 395 400

Gln Ala Glu Asp Asp Ala Asp Asp Xaa Phe Ser Pro Trp Gln Glu Gly
 405 410 415

Thr Asn Pro Xaa Leu Xaa Xaa Val Pro Asn Pro Val Phe Gly Ser Asp
 420 425 430

Thr Phe Cys Glu Pro Phe Asp Asp Ser Leu Leu Glu Glu Asp Phe Pro
 435 440 445

Asp Thr Gln Arg Ile Leu Thr Val Lys
 450 455

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<211> 193

<212> PRT

<213> Homo sapiens

<400> 6

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 1 5 10 15

Ala Ala Gly Val Tyr His Arg Ala Arg Ser Gly Lys Tyr Lys Thr Tyr
 20 25 30

Ala Ala Lys Ala Val Cys Gly Gly His Ala Thr Tyr Lys Ala Ala Arg
 35 40 45

Lys Gly His Val Cys Ala Ala Gly Trp Met Ala Lys Gly Arg Val Gly
 50 55 60

Tyr Val Lys Gly Asn Cys Gly Gly Lys Thr Gly Asp Tyr Gly Arg Asn
 65 70 75 80

Arg Ser Arg Trp Asp Ala Tyr Cys Tyr Asn His Ala Lys Cys Gly Gly
 85 90 95

Val Thr Asp Lys Arg Lys Ser Gly Asn Tyr Asp Asn Cys Tyr Trp His
 100 105 110

Arg Lys Tyr Gly Arg His Ser Asp Asp Asp Asp Gly Cys Ala Asp Tyr
 115 120 125

Val Tyr Asp Ser Tyr Asp Asp Val His Gly Val Gly Arg Tyr Cys Gly
 130 135 140

Asp Asp Asp Ser Thr Gly Asn Val Met Thr Lys Ser Asp Ala Ser Val
 145 150 155 160

Thr Ala Gly Gly Lys Tyr Val Ala Met Asp Val Ser Lys Ser Ser Gly
 165 170 175

Lys Asn Thr Ser Thr Thr Ser Thr Gly Asn Lys Asn Ala Gly Arg Ser
 180 185 190

His

<210> 7

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cctggggacg tttgccctgg ggccccagcc tggcccggt caccctggca tgaggag 117

atg ggc ctg ttg ctc ctg gtc cca ttg ctc ctg ctg ccc ggc tcc tac 165

Met Gly Leu Leu Leu Leu Val Pro Leu Leu Leu Leu Pro Gly Ser Tyr

1

5

10

15

gga ctg ccc ttc tac tac ggc ttc tac tac tcc aac agc gcc aac gac 213

Gly Leu Pro Phe Tyr Tyr Gly Phe Tyr Tyr Ser Asn Ser Ala Asn Asp

20

25

30

cag aac cta ggc aac ggt cat ggc aaa gac cta cnt aat gga gtg aag 261

Gln Asn Leu Gly Asn Gly His Gly Lys Asp Leu Xaa Asn Gly Val Lys

35

40

45

ctg gtg gtg gag aca ccc gag gag acc ctg ttc acc tac caa ggg gcc 309

Leu Val Val Glu Thr Pro Glu Glu Thr Leu Phe Thr Tyr Gln Gly Ala

50

55

60

agt gtg atc ctg ccc tgc cgc tac cgc tac gag ccg gcc ctg gtc tcc 357

Ser Val Ile Leu Pro Cys Arg Tyr Arg Tyr Glu Pro Ala Leu Val Ser

65

70

75

80

ccg cgg cgt gtg cgt gtc aaa tgg tgg aag ctg tcg gag aac ggg gcc	405
Pro Arg Arg Val Arg Val Lys Trp Trp Lys Leu Ser Glu Asn Gly Ala	
85 90 95	
cca gag aag gac gtg ctg gtg gcc atc ggg ctg agg cac cgc tcc ttt	453
Pro Glu Lys Asp Val Leu Val Ala Ile Gly Leu Arg His Arg Ser Phe	
100 105 110	
ggg gac tac caa ggc cgc gtg cac ctg cgg cag gac aaa gag cat gac	501
Gly Asp Tyr Gln Gly Arg Val His Leu Arg Gln Asp Lys Glu His Asp	
115 120 125	
gtc tcg ntg gag atc cag gnt ctg cgg ctg gag gac tat ggg cgt tac	549
Val Ser Xaa Glu Ile Gln Xaa Leu Arg Leu Glu Asp Tyr Gly Arg Tyr	
130 135 140	
cgc tgt gag gtc atn gac ggg ctg gag gat gaa agc ggt ctg gtg gag	597
Arg Cys Glu Val Xaa Asp Gly Leu Glu Asp Glu Ser Gly Leu Val Glu	
145 150 155 160	
ctg gag ctg cgg ggt gtg gtc ttt cct tac cag tcc ccc aac ggg cgc	645
Leu Glu Leu Arg Gly Val Val Phe Pro Tyr Gln Ser Pro Asn Gly Arg	
165 170 175	
tac cag ttc aac ttc cac gag ggc cag cag gtc tgt gca gag cag gct	693
Tyr Gln Phe Asn Phe His Glu Gly Gln Gln Val Cys Ala Glu Gln Ala	
180 185 190	
gcg gtg gtg gcc tcc ttt gag cag ctc ttc cgg gcc tgg gag gag ggc	741
Ala Val Val Ala Ser Phe Glu Gln Leu Phe Arg Ala Trp Glu Glu Gly	
195 200 205	
ctg gac tgg tgc aac gcg ggc tgg ctg cag gat gcc acg gtg cag tac	789
Leu Asp Trp Cys Asn Ala Gly Trp Leu Gln Asp Ala Thr Val Gln Tyr	
210 215 220	
ccc atc atg ttg ccc cgg cag ccc tgc ggt ggc ccg gac ctg gca cct	837
Pro Ile Met Leu Pro Arg Gln Pro Cys Gly Gly Pro Asp Leu Ala Pro	
225 230 235 240	

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Gly Leu Pro Phe Tyr Tyr Gly Phe Tyr Tyr Ser Asn Ser Ala Asn Asp
 20 25 30

Gln Asn Leu Gly Asn Gly His Gly Lys Asp Leu Xaa Asn Gly Val Lys
 35 40 45

Leu Val Val Glu Thr Pro Glu Glu Thr Leu Phe Thr Tyr Gln Gly Ala
 50 55 60

Ser Val Ile Leu Pro Cys Arg Tyr Arg Tyr Glu Pro Ala Leu Val Ser
 65 70 75 80

Pro Arg Arg Val Arg Val Lys Trp Trp Lys Leu Ser Glu Asn Gly Ala
85 90 95

Pro Glu Lys Asp Val Leu Val Ala Ile Gly Leu Arg His Arg Ser Phe
100 105 110

Gly Asp Tyr Gln Gly Arg Val His Leu Arg Gln Asp Lys Glu His Asp
115 120 125

Val Ser Xaa Glu Ile Gln Xaa Leu Arg Leu Glu Asp Tyr Gly Arg Tyr
130 135 140

Arg Cys Glu Val Xaa Asp Gly Leu Glu Asp Glu Ser Gly Leu Val Glu
145 150 155 160

Leu Glu Leu Arg Gly Val Val Phe Pro Tyr Gln Ser Pro Asn Gly Arg
165 170 175

Tyr Gln Phe Asn Phe His Glu Gly Gln Gln Val Cys Ala Glu Gln Ala
180 185 190

Ala Val Val Ala Ser Phe Glu Gln Leu Phe Arg Ala Trp Glu Glu Gly
195 200 205

Leu Asp Trp Cys Asn Ala Gly Trp Leu Gln Asp Ala Thr Val Gln Tyr
210 215 220

Pro Ile Met Leu Pro Arg Gln Pro Cys Gly Gly Pro Asp Leu Ala Pro
225 230 235 240

Gly Val Arg Ser Tyr Gly Pro Arg His Arg Arg Leu His Arg Tyr Asp
245 250 255

Val Phe Cys Phe Ala Thr Ala Leu Xaa Gly Arg Val Tyr Tyr Leu Xaa
260 265 270

His Pro Glu Xaa Leu Xaa Leu Thr Xaa Ala Arg Glu Ala Cys Gln Glu
275 280 285

Lys

<210> 9
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 <213> Homo sapiens

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Pro His Pro Asp Asn Ser Ser Leu Glu His Glu Arg Ile Ile His Ile
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Gln Glu Glu Asn Gly Pro Arg Leu Leu Val Val Ala Glu Gln Ala Lys
 35 40 45

Ile Phe Ser Gln Arg Gly Gly Asn Val Thr Leu Pro Cys Lys Phe Tyr
 50 55 60

His Glu His Thr Ser Thr Ala Gly Ser Gly Thr His Lys Ile Arg Val
 65 70 75 80

Lys Trp Thr Lys Leu Thr Ser Asp Tyr Leu Lys Glu Val Asp Val Phe
 85 90 95

Val Ala Met Gly His His Arg Lys Ser Tyr Gly Lys Tyr Gln Gly Arg
 100 105 110

Val Phe Leu Arg Glu Ser Ser Glu Asn Asp Ala Ser Leu Ile Ile Thr
 115 120 125

Asn Ile Met Leu Glu Asp Tyr Gly Arg Tyr Lys Cys Glu Val Ile Glu
 130 135 140

Gly Leu Glu Asp Asp Thr Ala Val Val Ala Leu Asn Leu Glu Gly Val
 145 150 155 160

Val Phe Pro Tyr Ser Pro Arg Leu Gly Arg Tyr Asn Leu Asn Phe His
 165 170 175

Glu Ala Gln Gln Ala Cys Leu Asp Gln Asp Ser Ile Ile Ala Ser Phe
 180 185 190

Asp Gln Leu Tyr Glu Ala Trp Arg Ser Gly Leu Asp Trp Cys Asn Ala
 195 200 205

Gly Trp Leu Ser Asp Gly Ser Val Gln Tyr Pro Ile Thr Lys Pro Arg
 210 215 220

Glu Pro Cys Gly Gly Lys Asn Thr Val Pro Gly Val Arg Asn Tyr Gly
 225 230 235 240

Phe Trp Asp Lys Glu Arg Ser Arg Tyr Asp Val Phe Cys Phe Thr Ser
 245 250 255

Asn Phe Asn Gly Arg Phe Tyr Tyr Leu Ile His Pro Thr Lys Leu Thr
 260 265 270

Tyr Asp Glu Ala Val Gln Ala Cys Leu Lys Asp Gly Ala Gln Ile Ala
 275 280 285

Lys Val Gly Gln Ile Phe Ala Ala Trp Lys Leu Leu Gly Tyr Asp Arg
 290 295 300

Cys Asp Ala Gly Trp Leu Ala Asp Gly Ser Val Arg Tyr Pro Ile Ser
 305 310 315 320

Arg Pro Arg Lys Arg Cys Ser Pro Asn Glu Ala Ala Val Arg Phe Val
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Gly Phe Pro Asp Lys Lys His Lys Leu Tyr Gly Val Tyr Cys Phe Arg
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Ala Tyr Asn
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 gacgggcaca gctgcacaga gatagacccc tgtgcagacg gccttaacgg aggggtgtcac 180
 gagcacgcca cctgtaag atg aca ggc ccg ggc aag cac aag tgt gag tgt 231
 Met Thr Gly Pro Gly Lys His Lys Cys Glu Cys
 1 5 10
 aaa agt cac tat gtc gga gat ggg ctg aac tgt gag ccg gag cag ctg 279
 Lys Ser His Tyr Val Gly Asp Gly Leu Asn Cys Glu Pro Glu Gln Leu
 15 20 25
 ccc att gac cgc tgc tta cag gac aat ggg cag tgc cat gca gac gcc 327
 Pro Ile Asp Arg Cys Leu Gln Asp Asn Gly Gln Cys His Ala Asp Ala
 30 35 40
 aaa tgt gtc gac ctc cac ttc cag gat acc act gtt ggg gtg ttc cat 375
 Lys Cys Val Asp Leu His Phe Gln Asp Thr Thr Val Gly Val Phe His
 45 50 55

cta cgc tcc cca ctg ggc cag tat aag ctg acc ttt gac aaa gcc aga	423
Leu Arg Ser Pro Leu Gly Gln Tyr Lys Leu Thr Phe Asp Lys Ala Arg	
60 65 70 75	
gag gcc tgt gcc aac gaa gct gcg acc atg gca acc tac aac cag ctc	471
Glu Ala Cys Ala Asn Glu Ala Ala Thr Met Ala Thr Tyr Asn Gln Leu	
80 85 90	
tcc tat nnc cag aag gcc aag tac cac ctg tgc tca gca ggc tgg ctg	519
Ser Tyr Xaa Gln Lys Ala Lys Tyr His Leu Cys Ser Ala Gly Trp Leu	
95 100 105	
gag acc ggg cgg gtt gcc tac ccc aca gcc ttc gcc tcc cag aac tgt	567
Glu Thr Gly Arg Val Ala Tyr Pro Thr Ala Phe Ala Ser Gln Asn Cys	
110 115 120	
ggc tct ggt gtg gtt ggg ata gtg gac tat gga cct aga ccc aac aag	615
Gly Ser Gly Val Val Gly Ile Val Asp Tyr Gly Pro Arg Pro Asn Lys	
125 130 135	
agt gaa atg tgg gat gtc ttc tgc tat cgg atg aaa gat gtg aac tgc	663
Ser Glu Met Trp Asp Val Phe Cys Tyr Arg Met Lys Asp Val Asn Cys	
140 145 150 155	
acc tnc aag gtg ggc tat gtg gga gat ggc ttc tca tac agt ggg aac	711
Thr Xaa Lys Val Gly Tyr Val Gly Asp Gly Phe Ser Tyr Ser Gly Asn	
160 165 170	
ctg ctg cag gtc ctg atg tcc ttc ccc tca ctc aca aac ttc ctg acg	759
Leu Leu Gln Val Leu Met Ser Phe Pro Ser Leu Thr Asn Phe Leu Thr	
175 180 185	
gaa gtg ctg gcc tat tcc aac agc tca gct cga ggc cgt gca ttt cta	807
Glu Val Leu Ala Tyr Ser Asn Ser Ser Ala Arg Gly Arg Ala Phe Leu	
190 195 200	
gaa cac ctg act gac ctg tcc atc cgc ggc acc ctc ttt gtn cca cag	855
Glu His Leu Thr Asp Leu Ser Ile Arg Gly Thr Leu Phe Val Pro Gln	
205 210 215	

aac agt ggg ctg ggg gag aat gag acc ttg tct ggg cgg gac atc gag	903
Asn Ser Gly Leu Gly Glu Asn Glu Thr Leu Ser Gly Arg Asp Ile Glu	
220 225 230 235	
 cac cac ctg gcc aat gtc agc atg ttt ttc tac aat gac ctt gtc aat	951
His His Leu Ala Asn Val Ser Met Phe Phe Tyr Asn Asp Leu Val Asn	
240 245 250	
 ggc acc acc ctg caa acg agg ctg gga agc aag ctg ctg atc act gac	999
Gly Thr Thr Leu Gln Thr Arg Leu Gly Ser Lys Leu Leu Ile Thr Asp	
255 260 265	
 aga cag gac cca ctg cac ccg acg gag acc agg tgt gtt gat gga aga	1047
Arg Gln Asp Pro Leu His Pro Thr Glu Thr Arg Cys Val Asp Gly Arg	
270 275 280	
 gac act ctg gag tgg gac atc tgt gcc tcc aat ggg atc aca cat gtc	1095
Asp Thr Leu Glu Trp Asp Ile Cys Ala Ser Asn Gly Ile Thr His Val	
285 290 295	
 att tcc agg yct tta aaa gca ccc cct gcc ccc gtg acc ttg ncc cac	1143
Ile Ser Arg Xaa Leu Lys Ala Pro Pro Ala Pro Val Thr Leu Xaa His	
300 305 310 315	
 act ggn ttg gga gna ggg atc ttc tnt gnc atc atc ctg gtg act ggg	1191
Thr Gly Leu Gly Xaa Gly Ile Phe Xaa Xaa Ile Ile Leu Val Thr Gly	
320 325 330	
 gct gtt gcc ttg gct gct tac tcc tac ttt cgg ata aac cgg aaa aca	1239
Ala Val Ala Leu Ala Ala Tyr Ser Tyr Phe Arg Ile Asn Arg Lys Thr	
335 340 345	
 atc ggc ttc can cat ttt ga	1259
Ile Gly Phe Xaa His Phe	
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<213> Homo sapiens

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			20					25					30		
Leu	Gln	Asp	Asn	Gly	Gln	Cys	His	Ala	Asp	Ala	Lys	Cys	Val	Asp	Leu
		35					40					45			
His	Phe	Gln	Asp	Thr	Thr	Val	Gly	Val	Phe	His	Leu	Arg	Ser	Pro	Leu
	50					55					60				
Gly	Gln	Tyr	Lys	Leu	Thr	Phe	Asp	Lys	Ala	Arg	Glu	Ala	Cys	Ala	Asn
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Glu	Ala	Ala	Thr	Met	Ala	Thr	Tyr	Asn	Gln	Leu	Ser	Tyr	Xaa	Gln	Lys
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Ala	Lys	Tyr	His	Leu	Cys	Ser	Ala	Gly	Trp	Leu	Glu	Thr	Gly	Arg	Val
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		115					120					125			
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	130					135					140				
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Tyr	Val	Gly	Asp	Gly	Phe	Ser	Tyr	Ser	Gly	Asn	Leu	Leu	Gln	Val	Leu
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		195					200					205			
Leu	Ser	Ile	Arg	Gly	Thr	Leu	Phe	Val	Pro	Gln	Asn	Ser	Gly	Leu	Gly
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Thr Arg Leu Gly Ser Lys Leu Leu Ile Thr Asp Arg Gln Asp Pro Leu
 260 265 270

His Pro Thr Glu Thr Arg Cys Val Asp Gly Arg Asp Thr Leu Glu Trp
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Asp Ile Cys Ala Ser Asn Gly Ile Thr His Val Ile Ser Arg Xaa Leu
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Lys Ala Pro Pro Ala Pro Val Thr Leu Xaa His Thr Gly Leu Gly Xaa
 305 310 315 320

Gly Ile Phe Xaa Xaa Ile Ile Leu Val Thr Gly Ala Val Ala Leu Ala
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Phe

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Gln Ala Ala Gly Val Tyr His Arg Glu Ala Arg Ala Gly Arg Tyr Lys
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Leu Thr Tyr Ala Glu Ala Lys Ala Val Cys Glu Phe Glu Gly Gly Arg
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Leu Ala Thr Tyr Lys Gln Leu Glu Ala Ala Arg Lys Ile Gly Phe His
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Val Cys Ala Ala Gly Trp Met Ala Lys Gly Arg Val Gly Tyr Pro Ile
 85 90 95

Val Lys Pro Gly Pro Asn Cys Gly Phe Gly Lys Thr Gly Ile Ile Asp
 100 105 110

Tyr Gly Ile Arg Leu Asn Arg Ser Glu Arg Trp Asp Ala Tyr Cys Tyr
 115 120 125

Asn Pro His Ala Lys Glu Cys Gly Gly Val Phe Thr Asp Pro Lys Arg
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Ile Phe Lys Ser Pro Gly Phe Pro Asn Glu Tyr Asp Asp Asn Gln Val
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Cys Tyr Trp His Ile Arg Leu Lys Tyr Gly Gln Arg Ile His Leu Ser
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Phe Leu Asp Phe Asp Leu Glu His Asp Pro Gly Cys Leu Ala Asp Tyr
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Val Glu Ile Tyr Asp Ser Tyr Asp Asp Val His Gly Phe Val Gly Arg
 195 200 205

Tyr Cys Gly Asp Glu Leu Pro Glu Asp Ile Ile Ser Thr Gly Asn Val
 210 215 220

Met Thr Leu Lys Phe Leu Ser Asp Ala Ser Val Thr Ala Gly Gly Phe
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Gln Ile Lys Tyr Val Thr Val Asp Pro Ala Ser Lys Ser Ser Gln Ala
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